

Amendments to the Specification:

Please replace the paragraph on page 24, lines 14-16, with the following amended paragraph:

The nucleotide sequence of siRNAs may be designed using a siRNA design computer program available from the Ambion website (http://www.ambion.com/techlib/misc/siRNA_finder.html). Nucleotide sequences for the siRNA are selected by the computer program based on the following protocol:

Please replace the paragraph on page 41, lines 23-32, with the following amended paragraph:

When gene-expression profiles of cancer cells from 18 pancreatic cancer patients using a cDNA microarray representing 23,040 human genes (Nakamura et al., 2003) were analyzed, 265 genes that were commonly up-regulated in pancreatic cancer cells were identified. Among them, the present inventors focused on a gene with in-house code C1958, which corresponded to EST Hs. 40530 in the UniGene database in NCBI (<http://www.ncbi.nlm.nih.gov/UniGene/>). Expression of this gene was elevated in 7 of 9 pancreatic cancers with greater signal intensities than the cut-off value on the microarray in comparison with normal pancreatic ducts (FIG. 1(a)). Subsequent semi-quantitative RT-PCR analysis confirmed elevated expression in 10 of 12 pancreatic cancers, and 2 of 5 pancreatic cancer cell line compared with normal pancreatic ducts (FIG. 1(b)).